# **SAXS**

#### The beamlines - current resources

#### Most common beamline elements in use

**Optics** 

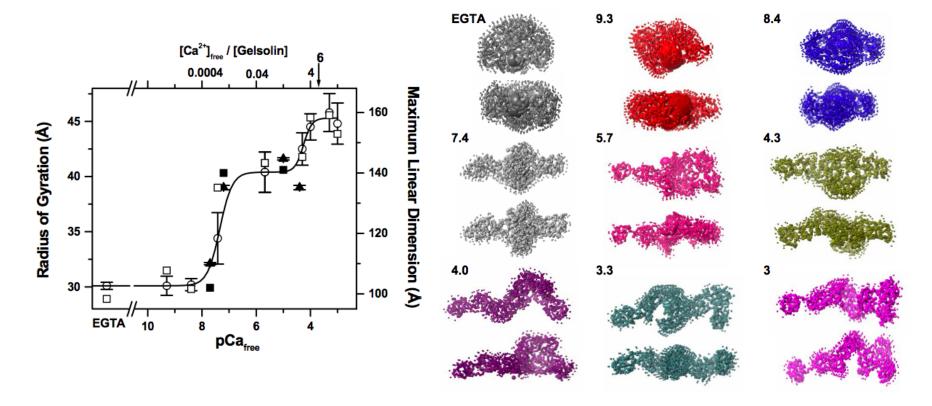
mono, focusing mirror, slits, evacuate beam path

**End-station** 

CCD, sample cell, syringe pump

## Research Program

Low resolution structural modeling of proteins in solution.



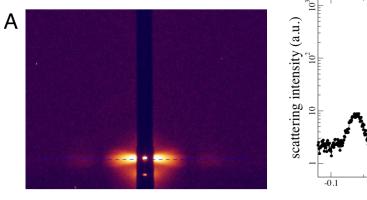
Conformation change of Gelsolin as a function of free Ca concentration Ashish et.al., J. Biol. Chem., 2007

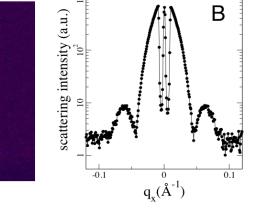
#### What can be done with a new source such as NSLSII

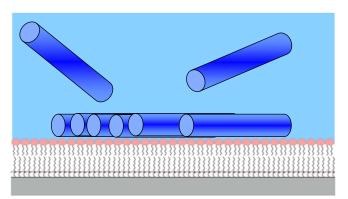
### **Unsolved Science Questions**

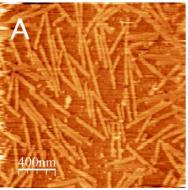
(Low resolution) molecular movies of protein-protein interaction. Membrane proteins.

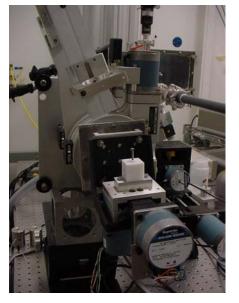
Source may not be the limiting factor.











Tobacco Mosaic Virus (TMV) adsorbed to a substrate submerged in a buffer solution

Yang, Wang, Fukuto, Cheeco on-going

### My next SAXS beamlines

#### **Beamlines**

ID beamline more challenging experiments that require high brightness BM beamline for protein solution scattering that will have high demand for capacity

### **Optics**

Not fundamentally different from NSLS beamlines

#### **End-station**

Better detectors: 2D pixel array detector, SAXS/WAXS Multiple probes for sample characterization (UV-VIS, CD, DLS, ...) Microfluidic sample cell that can perform solution processing Automatic sample handling

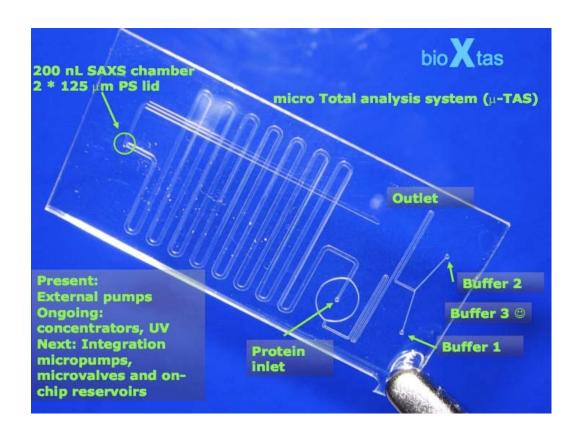
# Laboratories and ancillary facilities

#### Labs

Wet lab, protein purification facility

IT

Remote access





200nl sample volume, 0.5mm beam, 109ph/s, 5min exposure

On-going effort in Europe from Vestergaard presentation, University of Copenhagen

### How to we get ready for NSLS-II?

Instrumentation (detectors, microfluidics, ...)

Capability to support user research (staffing, supporting facilities, ...)

Methods for data analysis

Grow user community